SUMMARY OF THE INVENTION:

Calf-chymosin gene is isolated preferably from the fourth stomach of milk fed calf tissues. Recombinant calf-chymosin is produced by cloning chymosin gene with bacterial expression vector PET21b and is transformed to E-coli strain. This E-coli strain containing recombinant calf-chymosin gene is fermented under suitable conditions preferably in a culture medium developed by us. This medium contains the following

Peptone – 12g/l Yeast Extract – 24g/l Sodium chloride – 10g/l

Prochymosin produced during fermentation is subjected to denaturation by increasing the pH of the medium to 10-11. The suspension then diluted and the pH reduced to about 8 for effective renaturation of the protein. The prochymosin thus obtained is then acidified for activation and is further processed.

This invention relates to a process for producing recombinant calf-chymosin which comprises the steps of isolating calf-chymosin gene, cloning the same in bacterial expression vector PET 21b, transforming said cloned vector into cells of E-coli, fermenting said E-coli strains to produce pro-chymosin, converting said prochymosin to chymosin and subsequently recovering the recombinant calf chymosin.

This invention also includes recombinant calf-chymosin having the following amino acid sequence:

MetAlaSerIle ThrArgIle ProLeuTyr LysGlyLysSer LeuArgLys AlaLeuLys 1 ATGGCTAGCA TCACTAGGAT CCCTCTGTAC AAAGGCAAGT CTCTGAGGAA GGCGCTGAAG TACCGATCGT AGTGATCCTA GGGAGACATG TTTCCGTTCA GAGACTCCTT CCGCGACTTC GluHisGlyLeu LeuGluAsp PheLeuGln LysGlnGlnTyr GlyIleSer SerLysTyr 61 GAGCATGGGC TTCTGGAGGA CTTCCTGCAG AAACAGCAGT ATGGCATCAG CAGCAAGTAC CTCGTACCCG AAGACCTCCT GAAGGACGTC TTTGTCGTCA TACCGTAGTC GTCGTTCATG SerGlyPheGly GluValAla SerValPro LeuThrAsnTyr LeuAspSer GlnTyrPhe 121 TCCGGCTTCG GGGAGGTGGC CAGCGTGCCC CTGACCAACT ACCTGGATAG TCAGTACTTT AGGCCGAAGC CCCTCCACCG GTCGCACGGG GACTGGTTGA TGGACCTATC AGTCATGAAA GlyLysIleTyr LeuGlyThr ProProGln GluPheThrVal LeuPheAsp ThrGlySer 181 GGGAAGATCT ACCTCGGGAC CCCGCCCCAG GAGTTCACCG TGCTGTTTGA CACTGGCTCC CCCTTCTAGA TGGAGCCCTG GGGCGGGGTC CTCAAGTGGC ACGACAAACT GTGACCGAGG SerAspPheTrp ValProSer IleTyrCys LysSerAsnAla CysLysAsn HisGlnArg 241 TCTGACTTCT GGGTACCCTC TATCTACTGC AAGAGCAATG CCTGCAAAAA CCACCAGCGC AGACTGAAGA CCCATGGGAG ATAGATGACG TTCTCGTTAC GGACGTTTTT GGTGGTCGCG PheAspProArg LysSerSer ThrPheGln AsnLeuGlyLys ProLeuSer IleHisTyr 301 TTCGACCCGA GAAAGTCGTC CACCTTCCAG AACCTGGGCA AGCCCCTGTC TATCCACTAC AAGCTGGGCT CTTTCAGCAG GTGGAAGGTC TTGGACCCGT TCGGGGACAG ATAGGTGATG GlyThrGlyLys MetGlnGly IleLeuGly TyrAspThrVal ThrValSer AsnIleVal 361 GGGACAGGCA AGATGCAGGG GATCCTGGGC TATGACACCG TCACTGTCTC CAACATTGTG CCCTGTCCGT TCTACGTCCC CTAGGACCCG ATACTGTGGC AGTGACAGAG GTTGTAACAC AspIleGlnGln ThrValVal LeuSerThr GlnGluProGly AspValPhe ThrTyrAla 421 GACATCCAGC AGACAGTAGT CCTGAGCACC CAGGAGCCCG GGGACGTCTT CACCTATGCC CTGTAGGTCG TCTGTCATCA GGACTCGTGG GTCCTCGGGC CCCTGCAGAA GTGGATACGG GluPheAspGly IleLeuGly MetAlaTyr ProSerLeuAla SerGluVal LeuAspThr 481 GAATTCGACG GGATCCTGGG GATGGCGTAC CCCTCGCTGG CCTCAGAAGT ACTCGATACC CTTARGCTGC CCTAGGACCC CTACCGCATG GGGAGCGACC GGAGTCTTCA TGAGCTATGG 541 GGCTTTGACA ACATGATGAA CAGGCACCTG GTGGCCCAAG ACGTGTTCTC GGTTTACATG 601 GACAGGAATG GGCAGGGAAA CATGTTTACC CTGGGGGCCA TCGACCCGTC CTACTACACA 661 GGGTCCCTGC ACTGGGTGCC CGTGACAGTG CAGCAGTACT GGCAGTTCAC TGTGGACAGT 721 GTCACCATCA GCGGTGTGGT TGTGGCCTGT GAGGGTGGCT GTCAGGCCAT CCTGGACACG 781 GGCACCTCCA AGCTGGTCGG GCCCAGCAGC GACATCCTCA ACATCCAGCA GGCCATTGGA AlaThrGlnAsn GlnTyrAsp GluPheAsp IleAspCysAsp AsnLeuSer TyrMetPro 841 GCCACACAGA ACCAGTACGA TGAGTTTGAC ATCGACTGCG ACAACCTGAG CTACATGCCC 901 ACTGTGGTCT TTGAGATCAA TGGCAAAATG TACCCACTGA CCCCCTCCGC CTATACCAGC

GlyPheAspAsn MetMetAsn ArgHisLeu ValAlaGlnAsp ValPheSer ValTyrMet CCGAAACTGT TGTACTACTT GTCCGTGGAC CACCGGGTTC TGCACAAGAG CCAAATGTAC AspArgAsnGly GlnGlyAsn MetPheThr LeuGlyAlaIle AspProSer TyrTyrThr CTGTCCTTAC CCGTCCCTTT GTACAAATGG GACCCCCGGT AGCTGGGCAG GATGATGTGT GlySerLeuHis TrpValPro ValThrVal GlnGlnTyrTrp GlnPheThr ValAspSer CCCAGGGACG TGACCCACGG GCACTGTCAC GTCGTCATGA CCGTCAAGTG ACACCTGTCA ValThrIleSer GlyValVal ValAlaCys GluGlyGlyCys GlnAlaIle LeuAspThr CAGTGGTAGT CGCCACACCA ACACCGGACA CTCCCACCGA CAGTCCGGTA GGACCTGTGC GlyThrSerLys LeuValGly ProSerSer AspIleLeuAsn IleGlnGln AlaIleGly CCGTGGAGGT TCGACCAGCC CGGGTCGTCG CTGTAGGAGT TGTAGGTCGT CCGGTAACCT CGGTGTGTCT TGGTCATGCT ACTCAAACTG TAGCTGACGC TGTTGGACTC GATGTACGGG ThrValValPhe GluIleAsn GlyLysMet TyrProLeuThr ProSerAla TyrThrSer TGACACCAGA AACTCTAGTT ACCGTTTTAC ATGGGTGACT GGGGGAGGCG GATATGGTCG GlnAspGlnGly PheCysThr SerGlyPhe GlnSerGluAsn HisSerGln LysTrpIle 961 CAGGACCAGG GCTTCTGTAC CAGTGGCTTC CAGAGTGAAA ATCATTCCCA GAAATGGATC GTCCTGGTCC CGAAGACATG GTCACCGAAG GTCTCACTTT TAGTAAGGGT CTTTACCTAG LeuGlyAspVaT PhelleArg Glutyrtyr SerValPheAsp ArgalaAsn AsnLeuVal CTGGGGGATG TTTTCATCCG AGAGTATTAC AGCGTCTTTG ACAGGGCCAA CAACCTCGTG GACCCCCTAC AAAAGTAGGC TCTCATAATG TCGCAGAAAC TGTCCCGGTT GTTGGAGCAC

1081 GGGCTGGCCA AAGCCATCTG A CCCGACCGGT TTCGGTAGAC T

In the above sequence, amino acids shown in red indicate sequence variation of chymosin gene of our invention compared to the reported and published sequence.

TREPLACEMENT SHEET]

WO 2005/094185 PCT/IN2004/000074

PCR amplification of prepro chymosin was performed using the 50ng of 1st strand cDNA with a reverse primer (5'-TGT GGG GAC AGT GAG GTT CTT GGT C-3'), and a forward primer (5'-ATG AGG TGT CTC GTG GTG CTA CTT-3') in a thermal cycler programmed as (step 1: 95-5'; step 2: 94-30sec.; step 3: 54-30sec; step 4: 72-1min; step 5: go to step 2 34 times; step 6: 72-7min; step 7: end). The PCT reaction when analyzed on 1.0% agarose gel showed an amplified band of 1.2kb. The 1.2kb fragment was cut with a sterile blade and the gel slice was dissolved in 500µl of Tris saturated phenol was added and left in liquid nitrogen for a few min. The microcentrifuge tube was allowed to come to room temperature and centrifuged for 5min at 12,000rpm, 4°C. The upper aqueous phase was extracted with phenol: chloroform: isoamyl alcohol (25:24:1) and DNA was precipitated with 1/10th volume sodium acetate and 2.5 volume ethanol at -70°C for 1 h. DNA was precipitated at 15,000rpm for 15 min. The pellet was dried and dissolved in sterile distilled water. This eluted 1.2kb fragment was ligated at Smal site of pBSSK+ plasmid, which was then transformed in to TOP10 cells of E.coli. The recombinant clones were selected (blue white screening) and checked with restriction digestion analysis of the plasmids. Recombinant plasmid was taken as a template and a PCR was performed using a forward primer (5'-GAT ATA CAT ATG GCT AGC ATC ACT AGG ATC CCT CTG TAC-3') and reverse primer (5'-GCA GTA AGC TTG ACA GTG AGG TTC CTT GGT CAG CG-3') containing Nde 1 and Hind III sites. An amplified band of 1098bp was observed when the PCR product was analyzed on 1.0% agarose gel. This amplified fragment of 1098bp was eluted from the gel and ligated in pET21b expression vector at Nde 1 and Hind III sites and transformed in to BL21 cells of E.coli for the expression of the chymosin gene.

CLAIMS:

- 1. A process for producing recombinant calf-chymosin which comprises the steps of isolating calf-chymosin gene, cloning the same in bacterial expression vector PET21b, transforming said cloned vector into cells of E.coli, fermenting said E.coli strains to produce pro-chymosin, converting said pro-chymosin to chymosin and subsequently recovering the recombinant calf-chymosin.
- 2. The process as claimed in claim 1, wherein calf-chymosin gene is obtained by isolating RNA from the fourth stomach of calf tissue, synthesising a first strand of cDNA therefrom by treating the same with a reverse primer such as 5'-TGT GGG GAG AGT GAG GTT CTT GGT C-3' and then with a forward primer such as 5'-ATG AGG TGT CTC GTG GTG CTA CTT 3 and with a reverse primer such as 5'TGT GGT GAC AGT GAG GTT CTT GGT C-3'.
- 3. The process as claimed in claims 1 and 2 wherein said C DNA is ligated at small site of pBSSK+ plasmid and then transformed into TOP 10 cells of E-coli.
- 4. The process as claimed in claim 3 wherein said recombinant clones were identified and treated with a forward primer such as 5'-GAT ATA CAT ATG GCT AGC ATC ACT AGG ATC CCT CTG TAC 3' and reverse primer such as 5' GCA GTA AGC TTG ACA GTG TTC CTT GGT CAG CG-3' containing Nde I and Hind III sites to obtain an amplified fragment.
- 5. The process as claimed in claim 4 wherein said amplified fragment is transformed into cells of E.coli for expressing said chymosin gene.

WO 2005/094185 PCT/IN2004/000074

6. The process as claimed in any of the preceding claims wherein said E.coli cells containing recombinant calf chymosin gene is fermented in a medium containing 12g/L peptone, 24g/L of yeast extract and 10g/L of sodium chloride in the presence of supplements for fermentation and the suspended cells produced on completion of fermentation is lysed, chilled and pH adjusted to 8 before incubating at room temperature and the supernatent containing prochymosin is separated.

- 7. The process as claimed in claim 6, wherein the pH of said prochymosin containing supernatent is adjusted to 2 at room temperature and further incubated for about 6 hrs with gentle stirring and filtered.
- 8. The process as claimed in claim 7 wherein the pH of said filtrate is adjusted to about 5 and further incubated, filtered and treated with a solution containing sodium benzoate and thereafter a solution containing and sodium chloride to activate prochymosin to chymosin.
- 9. The process as claimed in claim 8 wherein the filtrate obtained after the addition of sodium benzoate solution is treated with a solution of sodium chloride under stirring and cooking, and the precipitate suspended in a chilled solution of 0.2M glycine with 0.001M EDTA and thereafter treated with 0.23% solution of sodium benzoate and stored under cooling.
- 10. The process as claimed in claim 9 wherein said chymosin obtained is formulated with 10% of sodium chloride and 0.2% of Trehalose.

Recombinant calf-chymosin having the following amino acid sequence: 11.

MetAlaSerIle ThrArgIle ProLeuTyr LysGlyLysSer LeuArgLys AlaLeuLys 361 GGGACAGGCA AGATGCAGGG GATCCTGGGC TATGACACCG TCACTGTCTC CAACATTGTG -541 GGCTTTGACA ACATGATGAA CAGGCACCTG GTGGCCCAAG ACGTGTTCTC GGTTTACATG > 721 GTCACCATCA GCGGTGTGGT TGTGGCCTGT GAGGGTGGCT GTCAGGCCAT CCTGGACACG 781 GGCACCTCCA AGCTGGTCGG GCCCAGCAGC GACATCCTCA ACATCCAGCA GGCCATTGGA -841 GCCACACAGA ACCAGTACGA TGAGTTTGAC ATCGACTGCG ACAACCTGAG CTACATGCCC 901 ACTGTGGTCT TTGAGATCAA TGGCAAAATG TACCCACTGA CCCCCTCCGC CTATACCAGC

1 ATGGCTAGCA TCACTAGGAT CCCTCTGTAC AAAGGCAAGT CTCTGAGGAA GGCGCTGAAG TACCGATCGT AGTGATCCTA GGGAGACATG TTTCCGTTCA GAGACTCCTT CCGCGACTTC GluHisGlyLeu LeuGluAsp PheLeuGln LysGlnGlnTyr GlyIleSer SerLysTyr 61 GAGCATGGGC TTCTGGAGGA CTTCCTGCAG AAACAGCAGT ATGGCATCAG CAGCAAGTAC CTCGTACCGG AAGACCTCCT GAAGGACGTC TTTGTCGTCA TACCGTAGTC GTCGTTCATG SerGlyPheGly GluValAla SerValPro LeuThrAsnTyr LeuAspSer GlnTyrPhe 121 TCCGGCTTCG GGGAGGTGGC CAGCGTGCCC CTGACCAACT ACCTGGATAG TCAGTACTTT AGGCCGAAGC CCCTCCACCG GTCGCACGGG GACTGGTTGA TGGACCTATC AGTCATGAAA GlyLysIleTyr LeuGlyThr ProProGln GluPheThrVal LeuPheAsp ThrGlySer 181 GGGAAGATCT ACCTCGGGAC CCCGCCCCAG GAGTTCACCG TGCTGTTTGA CACTGGCTCC CCCTTCTAGA TGGAGCCCTG GGGCGGGGTC CTCAAGTGGC ACGACAAACT GTGACCGAGG SerAspPheTrp ValProSer IleTyrCys LysSerAsnAla CysLysAsn HisGlnArg 241 TCTGACTTCT GGGTACCCTC TATCTACTGC AAGAGCAATG CCTGCAAAAA CCACCAGCGC AGACTGAAGA CCCATGGGAG ATAGATGACG TTCTCGTTAC GGACGTTTTT GGTGGTCGCG PheAspProArg LysSerSer ThrPheGln AsnLeuGlyLys ProLeuSer IleHisTyr 301 TTCGACCCGA GAAAGTCGTC CACCTTCCAG AACCTGGGCA AGCCCCTGTC TATCCACTAC AAGCTGGGCT CTTTCAGCAG GTGGAAGGTC TTGGACCCGT TCGGGGACAG ATAGGTGATG GlyThrGlyLys MetGlnGly IleLeuGly TyrAspThrVal ThrValSer AsnIleVal CCCTGTCCGT TCTACGTCCC CTAGGACCCG ATACTGTGGC AGTGACAGAG GTTGTAACAC AspIleGlnGln ThrValVal LeuSerThr GlnGluProGly AspValPhe ThrTyrAla 421 GACATCCAGC AGACAGTAGT CCTGAGCACC CAGGAGCCCG GGGACGTCTT CACCTATGCC CTGTAGGTCG TCTGTCATCA GGACTCGTGG GTCCTCGGGC CCCTGCAGAA GTGGATACGG GluPheAspGly IleLeuGly MetAlaTyr ProSerLeuAla SerGluVal LeuAspThr 481 GAATTCGACG GGATCCTGGG GATGGCGTAC CCCTCGCTGG CCTCAGAAGT ACTCGATACC -CTTAAGCTGC CCTAGGACCC CTACCGCATG GGGAGCGACC GGAGTCTTCA TGAGCTATGG GlyPheAspAsn MetMetAsn ArgHisLeu ValAlaGlnAsp ValPheSer ValTyrMet. CCGAAACTGT TGTACTACTT GTCCGTGGAC CACCGGGTTC TGCACAAGAG CCAAATGTAC AspArgAsnGly GlnGlyAsn MetPheThr LeuGlyAlaIle AspProSer TyrTyrThr-601 GACAGGAATG GGCAGGGAAA CATGTTTACC CTGGGGGCCA TCGACCCGTC CTACTACACA CTGTCCTTAC CCGTCCCTTT GTACAAATGG GACCCCCGGT AGCTGGGCAG GATGATGTGT GlySerLeuHis TrpValPro ValThrVal GlnGlnTyrTrp GlnPheThr ValAspSer 661 GGGTCCCTGC ACTGGGTGCC CGTGACAGTG CAGCAGTACT GGCAGTTCAC TGTGGACAGT — CCCAGGGACG TGACCCACGG GCACTGTCAC GTCGTCATGA CCGTCAAGTG ACACCTGTCA ValThrIleSer GlyValVal ValAlaCys GluGlyGlyCys GlnAlaIle LeuAspThr CAGTGGTAGT CGCCACACCA ACACCGGACA CTCCCACCGA CAGTCCGGTA GGACCTGTGC GlyThrSerLys LeuValGly ProSerSer AspIleLeuAsn IleGlnGln AlaIleGly CCGTGGAGGT TCGACCAGCC CGGGTCGTCG CTGTAGGAGT TGTAGGTCGT CCGGTAACCT AlaThrGlnAsn GlnTyrAsp GluPheAsp IleAspCysAsp AsnLeuSer TyrMetPro CGGTGTGTCT TGGTCATGCT ACTCAAACTG TAGCTGACGC TGTTGGACTC GATGTACGGG ThrValValPhe GluIleAsn GlyLysMet TyrProLeuThr ProSerAla TyrThrSer TGACACCAGA AACTCTAGTT ACCGTTTTAC ATGGGTGACT GGGGGAGGCG GATATGGTCG GlnAspGlnGly PheCysThr SerGlyPhe GlnSerGluAsn HisSerGln LysTrpIle

IREPLACEMENT SHEET?

- 961 CAGGACCAGG GCTTCTGTAC CAGTGGCTTC CAGAGTGAAA ATCATTCCCA GAAATGGATC GTCCTGGTCC CGAAGACATG GTCACCGAAG GTCTCACTTT TAGTAAGGGT CTTTACCTAG LeuGlyAspVal PheileArg Glutyrtyr SerValPheAsp ArgAlaAsn AsnLeuVal/ 3 60
- 1021 CTGGGGGATG TTTTCATCCG AGAGTATTAC AGCGTCTTTG ACAGGGCCAA CAACCTCGTG GACCCCCTAC AAAAGTAGGC TCTCATAATG TCGCAGAAAC TGTCCCGGTT GTTGGAGCAC GlyLeuAlaLys, Alaile***
- 1081 GGGCTGGCCA AAGCCATCTG A CCCGACCGGT TTCGGTAGAC T
- 13. Recombinant calf-chymosin when produced by a process according to any of the preceding claims.

[REPLACEMENT SHEET]

ABSTRACT

Calf-chymosin containing 1098 p coding for 366 amino acids was isolated and cloned into a bacterial expression vector pET 21b. This strain is fermented and the protein is precipitated by addition of alkali. Active form is recovered by renaturation and purified under low pH.

Chymosin.ST25 SEQUENCE LISTING

Sudershan Biotech Limited Sudershan Biotech Limited <120> Recombinant Calf Chymosin and a process for producing the same <130> PCT0602 <140> PCT/IN04/000074 <141> 2004-03-30 <160> <170> PatentIn version 3.3 <210> 366 <211> <212> PRT <213> calf <400> Met Ala Ser Ile Thr Arg Ile Pro Leu Tyr Lys Gly Lys Ser Leu Arg
1 10 15 Lys Ala Leu Lys Glu His Gly Leu Leu Glu Asp Phe Leu Gln Lys Gln
20 25 30 Gln Tyr Gly Ile Ser Ser Lys Tyr Ser Gly Phe Gly Glu Val Ala Ser 35 40 45 Val Pro Leu Thr Asn Tyr Leu Asp Ser Gln Tyr Phe Gly Lys Ile Tyr 50 55 60 Leu Gly Thr Pro Pro Gln Glu Phe Thr Val Leu Phe Asp Thr Gly Ser 65 70 75 80 Ser Asp Phe Trp Val Pro Ser Ile Tyr Cys Lys Ser Asn Ala Cys Lys 85 90 95 Asn His Gln Arg Phe Asp Pro Arg Lys Ser Ser Thr Phe Gln Asn Leu 100 105 110

Gly Lys Pro Leu Ser Ile His Tyr Gly Thr Gly Lys Met Gln Gly Ile 115 120 125

Leu Gly Tyr Asp Thr Val Thr Val Ser Asn Ile Val Asp Ile Gln Gln 130 135 140

Thr Val Val Leu Ser Thr Gln Glu Pro Gly Asp Val Phe Thr Tyr Ala 145 150 155 160

115

Page 1

[REPLACEMENT SHEET]

Glu Phe Asp Gly Ile Leu Gly Met Ala Tyr Pro Ser Leu Ala Ser Glu 165 170 175 Val Leu Asp Thr Gly Phe Asp Asn Met Met Asn Arg His Leu Val Ala Gln Asp Val Phe Ser Val Tyr Met Asp Arg Asn Gly Gln Gly Asn Met 195 200 205 Phe Thr Leu Gly Ala Ile Asp Pro Ser Tyr Tyr Thr Gly Ser Leu His 210 215 220 Trp Val Pro Val Thr Val Gln Gln Tyr Trp Gln Phe Thr Val Asp Ser 225 230 235 240 Val Thr Ile Ser Gly Val Val Ala Cys Glu Gly Gly Cys Gln Ala 245 250 255 Ile Leu Asp Thr Gly Thr Ser Lys Leu Val Gly Pro Ser Ser Asp Ile 260 265 270 Leu Asn Ile Gln Gln Ala Ile Gly Ala Thr Gln Asn Gln Tyr Asp Glu 275 280 285 Phe Asp Ile Asp Cys Asp Asn Leu Ser Tyr Met Pro Thr Val Val Phe 290 300 Glu Ile Asn Gly Lys Met Tyr Pro Leu Thr Pro Ser Ala Tyr Thr Ser 305 310 315 320 Gln Asp Gln Gly Phe Cys Thr Ser Gly Phe Gln Ser Glu Asn His Ser 325 330 335 Gln Lys Trp Ile Leu Gly Asp Val Phe Ile Arg Glu Tyr Tyr Ser Val 340 345 350 Phe Asp Arg Ala Asn Asn Leu Val Gly Leu Ala Lys Ala Ile 355 360 365 1101 DNA calf <400> atggctagca tcactaggat ccctctgtac aaaggcaagt ctctgaggaa ggcgctgaag gagcatgggc ttctggagga cttcctgcag aaacagcagt atggcatcag cagcaagtac

tccggcttcg gggaggtggc cagcgtgccc ctgaccaact acctggatag tcagtacttt

[REPLACEMENT SHEET]

Page 2

60

120 180

Chymosin.ST25

gggaag	atct acctcgggac	cccgccccag	gagttcaccg	tgctgtttga	cactggctcc	240
tctgac	ttct gggtaccctc	tatctactgc	aagagcaatg	cctgcaaaaa	ccaccagcgc	300
ttcgac	ccga gaaagtcgtc	caccttccag	aacctgggca	agcccctgtc	tatccactac	360
gggaca	ggca agatgcaggg	gatcctgggc	tatgacaccg	tcactgtctc	caacattgtg	420
gacatc	cagc agacagtagt	cctgagcacc	caggagcccg	gggacgtctt	cacctatgcc	480
gaattc	gacg ggatcctggg	gatggcgtac	ccctcgctgg	cctcagaagt	actcgatacc	540
ggcttt	gaca acatgatgaa	caggcacctg	gtggcccaag	acgtgttctc	ggtttacatg	600
gacagg	aatg ggcagggaaa	catgtttacc	ctgggggcca	tcgacccgtc	ctactacaca	660
gggtcc	ctgc actgggtgcc	cgtgacagtg	cagcagtact	ggcagttcac	tgtggacagt	720
gtcacc	atca gcggtgtggt	tgtggcctgt	gagggtggct	gtcaggccat	cctggacacg	780
ggcacc	tcca agctggtcgg	gcccagcagc	gacatcctca	acatccagca	ggccattgga	840
gccaca	caga accagtacga	tgagtttgac	atcgactgcg	acaacctgag	ctacatgccc	900
actgtg	gtct ttgagatcaa	tggcaaaatg	tacccactga	cccctccgc	ctataccagc	960
caggaccagg gcttctgtac cagtggcttc cagagtgaaa atcattccca gaaatggatc					1020	
ctgggggatg ttttcatccg agagtattac agcgtctttg acagggccaa caacctcgtg					1080	
gggctg	gcca aagccatctg	a		•		1101
<210> 3 <211> 25 <212> DNA <213> Artificial sequence <220> <223> Primer (reverse) for amplification of prepro chymosin						
<400> 3 tgtggggaca gtgaggttct tggtc					25	
<210> <211> <212> <213>	4 24 DNA Artificial sequ	uence				
<220> <223> Primer (forward) for amplification of prepro chymosin						
<400> 4 atgaggtgtc tcgtggtgct actt 24						
<210> <211> <212> <213>	5 39 DNA Artificial sequ	ience				

Page 3

220	Chymosin.ST25			
<220> <223>	Primer (forward) for amplification of recombinant plasmid			
	5 cata tggctagcat cactaggatc cctctgtac	39		
<210> <211> <212> <213>	6 35 DNA Artificial sequence			
<220> <223>	Primer (reverse) containing Nde 1 and Hind III sites			
	6 agct tgacagtgag gttccttggt cagcg	35		